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## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

- (i) APPLICANTS: Allaway, Graham P Dragic, Tatjana Litwin, Virginia M Maddon, Paul J Moore, John P
- (ii) TITLE OF INVENTION: Uses Of A Chemokine Receptor For Inhibiting HIV-1 Infection
- (iii) NUMBER OF SEQUENCES: 30
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: Cooper & Dunham LLP
    - (B) STREET: 1185 Avenue of the Americas
    - (C) CITY: New York
    - (D) STATE: New York
    - (E) COUNTRY: U.S.A.
  - (F) ZIP: 10036
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER:
  - (B) FILING DATE:
  - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:

  - (A) NAME: White, John P
    (B) REGISTRATION NUMBER: 28,678
  - (C) REFERENCE/DOCKET NUMBER: 51320
  - (ix) TELECOMMUNICATION INFORMATION:
    - (A) TELEPHONE: 212-278-0400
    - (B) TELEFAX: 212-391-0525
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 38 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA (genomic)
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

CAAGGCTACT TCCCTGATTG GCAGAACTAC ACACCAGG

(2) INFORMATION FOR SEQ ID NO:2:

38

(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 25 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:	
AGCAAGCC	GA GTCCTGCGTC GAGAG	25
(2) INFO	RMATION FOR SEQ ID NO:3:	
<b>(i)</b>	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:3:	
	CC GCTGGGGACT TTC	23
	RMATION FOR SEQ ID NO:4:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:4:	
CCTGTTCG	GG CGCCACTGCT AGAGATTTTC CAC	33
(2) INFO	DRMATION FOR SEQ ID NO:5:	
(1)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 31 amino acids  (B) TYPE: amino acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: peptide	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:	
Met 1	Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr 5 10 15	
Set	Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Arg 20 25 30	-
(2) INFO	DRMATION FOR SEQ ID NO:6:	
/ 2 \	CROTTENCE CHARACTERISTICS:	

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- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

His Tyr Ala Ala Ala Gln Trp Asp Phe Gly Asn Thr Met Cys Gln

- (2) INFORMATION FOR SEQ ID NO:7:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 32 amino acids (B) TYPE: amino acid

    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

Arg Ser Gln Lys Glu Gly Leu His Tyr Thr Cys Ser Ser His Phe Pro

Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys Ile Val

- (2) INFORMATION FOR SEQ ID NO:8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 17 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser Asn Arg Leu Asp

Gln

- (2) INFORMATION FOR SEQ ID NO:9:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 36 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA (genomic)
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

AAGCTTGGAG AACCAGCGGT TACCATGGAG GGGATC	36
(2) INFORMATION FOR SEQ ID NO:10:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	30
GTCTGAGTCT GAGTCAAGCT TGGAGAACCA	30
(2) INFORMATION FOR SEQ ID NO:11:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 41 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	41
CTCGAGCATC TGTGTTAGCT GGAGTGAAAA CTTGAAGACT C	41
(2) INFORMATION FOR SEQ ID NO:12:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	30
GTCTGAGTCT GAGTCCTCGA GCATCTGTGT	30
(2) INFORMATION FOR SEQ ID NO:13:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 32 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	32
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(2) INFORMATION FOR SEQ ID NO:14:

		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:14:	
.;	GTC	TGAGT	CT GAGTCAAGCT TCAGAGAGAA	30
	(2)	INFO	RMATION FOR SEQ ID NO:15:	
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 32 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:15:	
	CTC	GAGCT	GA GTCAGAACCC AGCAGAGAGT TC	32
	(2)	INFO	RMATION FOR SEQ ID NO:16:	
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:16:	
	GTC:	rgagt	CT GAGTCCTCGA GCTGAGTCAG	30
	(2)	INFO	RMATION FOR SEQ ID NO:17:	
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 32 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
		(ii)	MOLECULE TYPE: DNA (genomic)	
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:17:	
	AAGO	CTTCAC	T ACATCCACAA CATGCTGTCC AC	32
	(2)	INFO	RMATION FOR SEQ ID NO:18:	
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	



(ii) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

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	(ii) MOLECULE TYPE: DNA (genomic)		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:		
GTCT	TGAGTCT GAGTCAAGCT TCAGTACATC	30	
(2)	INFORMATION FOR SEQ ID NO:19:		
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 31 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
	(ii) MOLECULE TYPE: DNA (genomic)		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:		
CTCG	BAGCCTC GTTTTATAAA CCAGCCGAGA C	31	
(2)	INFORMATION FOR SEQ ID NO:20:		
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
	(ii) MOLECULE TYPE: DNA (genomic)		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:		
GTCT	GAGTCT GAGTCCTCGA GCCTCGTTTT	30	
(2)	) INFORMATION FOR SEQ ID NO:21:		
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 29 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
	(ii) MOLECULE TYPE: DNA (genomic)		
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:		
AAGC	TTCAGG GAGAAGTGAA ATGACAACC	29	
(2)	INFORMATION FOR SEQ ID NO:22:		
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		

	GTCI	rgagtct gagtcaagct tcagggagaa	30	
	(2)	INFORMATION FOR SEQ ID NO:23:		
		<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 33 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>		
e d	1	(ii) MOLECULE TYPE: DNA (genomic)		
	••	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:		
	CTCG	BAGCAGA CCTAAAACAC AATAGAGAGT TCC	33	
	(2)	INFORMATION FOR SEQ ID NO:24:		
		<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 30 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>		
		(ii) MOLECULE TYPE: DNA (genomic)		
		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:		
	GTCTGAGTCT GAGTCCTCGA GCAGACCTAA 30			
	(2) INFORMATION FOR SEQ ID NO:25:			
		(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 34 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii) MOLECULE TYPE: DNA (genomic)		
		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:		
	AAGO	CTTCTGT AGAGTTAAAA AATGAACCCC ACGG	34	
	(2) INFORMATION FOR SEQ ID NO:26:			
		<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 30 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>		
		(ii) MOLECULE TYPE: DNA (genomic)		
		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:		
	GTCT	TGAGTCT GAGTCAAGCT TCTGTAGAGT	30	
	(2)	INFORMATION FOR SEQ ID NO:27:		





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	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 34 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:27:	
CTC	GAGCC	AT TTCATTTTC TACAGGACAG CATC	34
(2)	INFO	RMATION FOR SEQ ID NO:28:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:28:	
GTC'	rgagt(	CT GAGTCCTCGA GCCATTTCAT	30
(2)	INFO	RMATION FOR SEQ ID NO:29:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:29:	
GTC:	TGAGT	CT GAGTCAAGCT TAACAAGATG GATTATCAA	39
(2)	INFO	RMATION FOR SEQ ID NO:30:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:30:	
GTCT	GAGT	CT GAGTCCTCGA GTCCGTGTCG CAAGCCCAC	39